

# PHGenFit: Path length extraction

Previous interface:

```
genfit::MeasuredStateOnPlane* Track::extrapolateToXXX(... , const int tr_point_id = 0)
```

Added a new interface that returns the path length, old interface also still available.

```
double Track::extrapolateToXXX(genfit::MeasuredStateOnPlane& state, ... , const int tr_point_id = 0)
```

The path length is calculated w.r.t. the given measurement (default 0).

Merged into J. Lajoie's repository, would come to the sPHENIX repository with [P.R. #201](#)

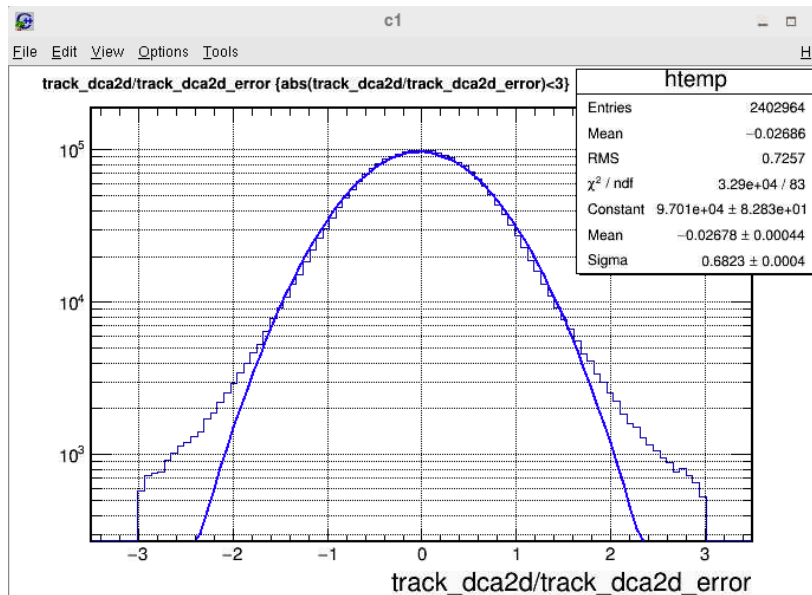
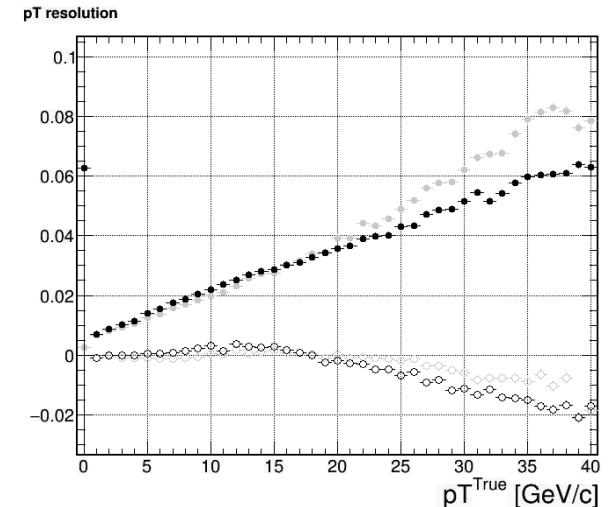
- <https://github.com/johnlajoie/coresoftware/pull/6>

# Investigate momentum/dca reconstruction Ladder MAPS + TPC

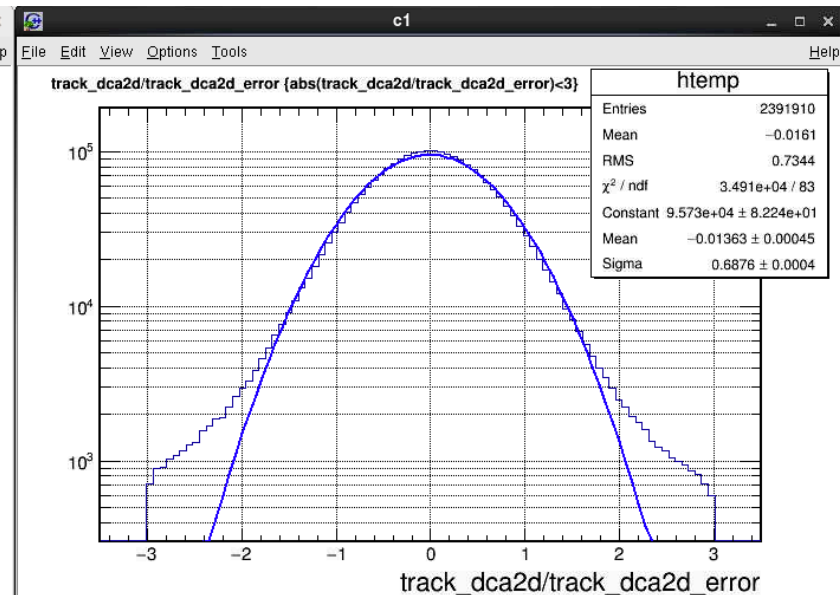
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# Problems

- ❖ pT offset is off 0
- ❖ dca2d pull  $\sigma$  ( $\sigma(\text{dca2d}/\text{error})$ ) is off 1.
- Possible origins:
  - field map inconsistency
  - clustering (center values, uncertainties)
- I checked the field map settings, **and it seems I used consistent field maps:**
  - /phenix/upgrades/decadal/fieldmaps/sPHENIX.2d.root
  - Scaled by 1.4/1.5
- We have a FastSim (truth tracking) module "PHG4TrackFastSim" which can do Kalman Fitting based on smeared PHG4Hits.
  - The central value is the PHG4Hit average position of the in and out hits
  - The smearing is given according to detector resolution
  - So there is no clustering involved



Alan

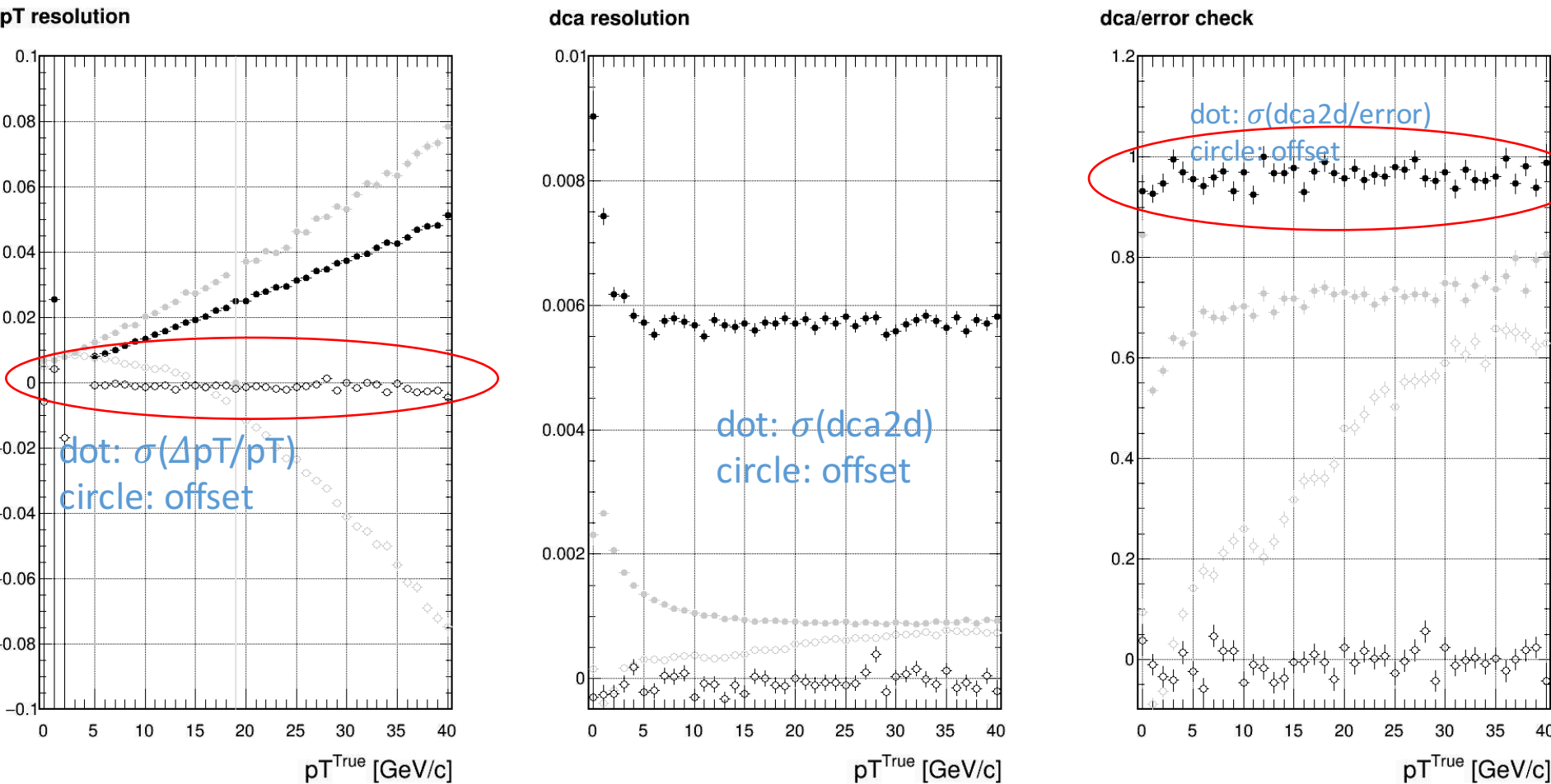


GenFit

# Alan (no PHG4SvtxMomentumRecal, grey) vs. FastSim/Truth Tracking (Black)

## FastSim Setup:

- Takes both MAPS and TPC PHG4Hits
- Assign 100 micron resolution to both phi and z resolution
  - All the absolute resolutions don't reflect true detector performance
  - Offsets and pull distributions still make sense
- The reconstructed  $p_T$  offset is 0.
- The dca2d pull  $\sigma$  is close to 1.



# Alan (no PHG4SvtxMomentumRecal, grey) vs. GenFit (with correct norm vectors, Black)

- GenFit refitted results:
  - Better pT resolution at high pT
  - Better dca2d resolution at low pT
  - Dca2d offsets are closer to 0

